

SEQUENCE LISTING

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<120> RECOMBINANT FUSOBACTERIUM NECROPHORUM LEUKOTOXIN VACCINE AND PREPARATION THEREOF

<130> PCT-30296

<160> 15

<170> PatentIn version 3.0

<210> 1

<211> 3241

<212> PRT

<213> Fusobacterium necrophorum

<400> 1

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Lys Ile Glu Lys Lys Asp Asn Val Tyr Asp Ile Thr Thr Asn Lys Ile
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Gln Gly Glu Asn Ala Phe Asn Ser Phe Asn Arg Phe Ala Leu Thr Glu
65      70      75      80
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Asn Asn Leu Phe Asn Phe Val Asn Gly Lys Ile Glu Val Asp Gly Ile
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Ser Phe His Ser Ile Ile Pro Lys Gln Asp Asp Phe Lys Lys Ala Leu
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Glu Glu Ala Lys His Gly Lys Val Phe Asn Gly Ile Ile Pro Val Asp
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Gly Lys Val Lys Ile Pro Leu Asn Pro Asn Gly Ser Ile Thr Val Glu
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Gly Lys Ile Asn Ala Val Glu Gly Ile Gly Leu Tyr Ala Ala Asp Ile
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Leu Lys Ala Thr Lys Thr Lys Ser Gly Asp Ile Ile Leu Ser Ala His
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Asp Ala Val Leu Glu Ala Asp Gly Asn Ile Lys Ile Ser Ala Lys Ala
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Thr Asn Gly Arg Phe Ile Lys Lys Glu Gly Glu Lys Glu Thr Tyr Asn
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Gly Lys Val Ile Gly Lys Asn Val Asp Ile Thr Ala Glu Ala Lys Asn
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Ser Ala Lys Ser Asn Ser Asn Val Thr Ile Glu Gly Glu Val Lys Ser
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Lys Gly Arg Ala Asp Ile Thr Ser Lys Ser Glu Asn Thr Ile Asp Ala

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 Thr Asp Ser Met Asp Ser Leu Thr Glu Lys Ile Thr Asn Tyr Ile
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Ser Ala 1250	Gly Leu Ala Gly 1255	Ala Ala Val Asn 1260	Asn Ile Gln Ser 1260
Lys Thr 1265	Ser Ala Leu Val 1270	Asn Ser Asp Ile 1275	Arg Asn Ala Asn 1275
Lys Phe 1280	Lys Val Asn Ala 1285	Ser Gly Gly Thr 1290	Gln Val Ala Ala 1290
Gly Ala 1295	Gly Leu Glu Ala 1300	Lys Glu Ser Gly 1305	Gly Gln Gly Lys 1305
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Glu Val 1325	Ser Ala Lys Ser 1330	Asn Asn Thr Val 1335	Ala Gly Glu Ser 1335
Glu Ser 1340	Gln Lys Met Asp 1345	Asp Val Thr Ala 1350	Tyr Gln Ala Asp 1350
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Val Asn 1385	Ala Ser Ile Ser 1390	Gly Arg Tyr Thr 1395	Asn Val Asn Arg 1395
Ala Asp 1400	Ala Lys Ala Leu 1405	Ala Thr Thr Gln 1410	Val Thr Ala Ala 1410
Val Thr 1415	Thr Gly Gly Thr 1420	Ser Ser Gly Ala 1425	Gly Leu Gly Asn 1425
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Ala Ser 1445	Val Asp Lys Ser 1450	Ile Glu Gly Ala 1455	Asn Glu Ile Asn 1455
Val Ile 1460	Ala Lys Asp Val 1465	Gly Ser Ser Asp 1470	Leu Ala Lys Glu 1470
Tyr Gln 1475	Ala Leu Leu Asn 1480	Lys Asp Lys Lys 1485	Tyr Leu Glu Asp 1485
Arg Gly 1490	Ile Asn Thr Thr 1495	Asn Gly Tyr Tyr 1500	Thr Lys Glu Gln 1500
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Val	Lys	Asn	Ser	Lys	Ile	Met	Thr	Arg	Asn	Lys	Leu	Asp	Val	Ile
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Arg	Val	His	Asn	Asn	Ser	Asp	Leu	Thr	Ser	Val	Arg	Glu	Lys	Val
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Gly	Asn	Val	Ser	Gln	Leu	Arg	Ala	Lys	Ala	Leu	Ala	Lys	Asn	Glu
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Asn	Val	Asp	Val	Asn	Asn	Val	Ile	Lys	Lys	Asn	Val	Glu	Ala	Lys
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Gln	Ala	Phe	Thr	Arg	Ala	Lys	Val	Asn	Ile	Leu	Gly	Lys	Gly	Asp
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Ala	Ala	Ala	Ala	Ala	Ala	Ile	Ser	Asn	Val	His	Ile	Ser	Asn	Glu
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3185						3190					3195			
Ala	Asn	Lys	Lys	Thr	Ser	Val	Gly	Ser	Asn	Gln	Val	Asp	Trp	Tyr
3200						3205					3210			
Thr	Asp	Lys	Tyr	Thr	Trp	His	Ser	Ser	Glu	Lys	Ala	Tyr	Lys	Lys
3215						3220					3225			

PCT302~1

Leu Thr Tyr Gln Ser Lys Arg Gly Glu Lys Gly Lys Lys
 3230 3235 3240

<210> 2

<211> 369

<212> PRT

<213> Fusobacterium necrophorum

<400> 2

Met Ser Gly Ile Lys Asn Asn Val Gln Arg Thr Arg Lys Arg Ile Ser
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 20 25 30
 Thr Val Arg Ala Asn Asp Thr Ile Thr Ala Thr Glu Asn Phe Gly Thr
 35 40 45
 Lys Ile Glu Lys Lys Asp Asn Val Tyr Asp Ile Thr Thr Asn Lys Ile
 50 55 60
 Gln Gly Glu Asn Ala Phe Asn Ser Phe Asn Arg Phe Ala Leu Thr Glu
 65 70 75 80
 Asn Asn Ile Ala Asn Leu Tyr Phe Gly Glu Lys Asn Ser Thr Gly Val
 85 90 95
 Asn Asn Leu Phe Asn Phe Val Asn Gly Lys Ile Glu Val Asp Gly Ile
 100 105 110
 Ile Asn Gly Ile Arg Glu Asn Lys Ile Gly Gly Asn Leu Tyr Phe Leu
 115 120 125
 Ser Ser Glu Gly Met Ala Val Gly Lys Asn Gly Val Ile Asn Ala Gly
 130 135 140
 Ser Phe His Ser Ile Ile Pro Lys Gln Asp Asp Phe Lys Lys Ala Leu
 145 150 155 160
 Glu Glu Ala Lys His Gly Lys Val Phe Asn Gly Ile Ile Pro Val Asp
 165 170 175
 Gly Lys Val Lys Ile Pro Leu Asn Pro Asn Gly Ser Ile Thr Val Glu
 180 185 190
 Gly Lys Ile Asn Ala Val Glu Gly Ile Gly Leu Tyr Ala Ala Asp Ile
 195 200 205
 Arg Leu Lys Asp Thr Ala Ile Leu Lys Thr Gly Ile Thr Asp Phe Lys
 210 215 220
 Asn Leu Val Asn Ile Ser Asp Arg Ile Asn Ser Gly Leu Thr Gly Asp
 225 230 235 240
 Leu Lys Ala Thr Lys Thr Lys Ser Gly Asp Ile Ile Leu Ser Ala His
 245 250 255
 Ile Asp Ser Pro Gln Lys Ala Met Gly Lys Asn Ser Thr Val Gly Lys
 260 265 270
 Arg Ile Glu Glu Tyr Val Lys Gly Asn Thr Lys Ala Asn Ile Glu Ser

PCT302~1

275		280		285											
Asp	Ala	Val	Leu	Glu	Ala	Asp	Gly	Asn	Ile	Lys	Ile	Ser	Ala	Lys	Ala
	290					295					300				
Thr	Asn	Gly	Arg	Phe	Ile	Lys	Lys	Glu	Gly	Glu	Lys	Glu	Thr	Tyr	Asn
	305				310					315					320
Thr	Pro	Leu	Ser	Leu	Ser	Asp	Val	Glu	Ala	Ser	Val	Arg	Val	Asn	Lys
				325					330					335	
Gly	Lys	Val	Ile	Gly	Lys	Asn	Val	Asp	Ile	Thr	Ala	Glu	Ala	Lys	Asn
			340					345					350		
Phe	Tyr	Asp	Ala	Thr	Leu	Val	Thr	Lys	Leu	Ala	Lys	His	Ser	Phe	Ser
		355					360					365			

Phe

<210> 3

<211> 927

<212> PRT

<213> Fusobacterium necrophorum

<400> 3

Gly	Arg	Phe	Ile	Lys	Lys	Glu	Gly	Glu	Lys	Glu	Thr	Tyr	Asn	Thr	Pro
1				5					10					15	
Leu	Ser	Leu	Ser	Asp	Val	Glu	Ala	Ser	Val	Arg	Val	Asn	Lys	Gly	Lys
			20					25					30		
Val	Ile	Gly	Lys	Asn	Val	Asp	Ile	Thr	Ala	Glu	Ala	Lys	Asn	Phe	Tyr
		35				40						45			
Asp	Ala	Thr	Leu	Val	Thr	Lys	Leu	Ala	Lys	His	Ser	Phe	Ser	Phe	Val
	50					55					60				
Thr	Gly	Ser	Ile	Ser	Pro	Ile	Asn	Leu	Asn	Gly	Phe	Leu	Gly	Leu	Leu
	65				70					75					80
Thr	Ser	Lys	Ser	Ser	Val	Val	Ile	Gly	Lys	Asp	Ala	Lys	Val	Glu	Ala
				85					90					95	
Thr	Glu	Gly	Lys	Ala	Asn	Ile	His	Ser	Tyr	Ser	Gly	Val	Arg	Ala	Thr
			100					105					110		
Met	Gly	Ala	Ala	Thr	Ser	Pro	Leu	Lys	Ile	Thr	Asn	Leu	Tyr	Leu	Glu
		115					120					125			
Lys	Ala	Asn	Gly	Lys	Leu	Leu	Ser	Ile	Gly	Ala	Gly	Tyr	Ile	Ser	Ala
		130				135					140				
Lys	Ser	Asn	Ser	Asn	Val	Thr	Ile	Glu	Gly	Glu	Val	Lys	Ser	Lys	Gly
					150					155					160
Arg	Ala	Asp	Ile	Thr	Ser	Lys	Ser	Glu	Asn	Thr	Ile	Asp	Ala	Ser	Val
				165					170					175	
Ser	Val	Gly	Thr	Met	Arg	Asp	Ser	Asn	Lys	Val	Ala	Leu	Ser	Val	Leu
			180					185					190		

PCT302~1

Val	Thr	Glu	Gly	Glu	Asn	Lys	Ser	Ser	Val	Lys	Ile	Ala	Lys	Gly	Ala
		195					200					205			
Lys	Val	Glu	Ser	Glu	Thr	Asp	Asp	Val	Asn	Val	Arg	Ser	Glu	Ala	Ile
	210					215					220				
Asn	Ser	Ile	Arg	Ala	Ala	Val	Lys	Gly	Gly	Leu	Gly	Asp	Ser	Gly	Asn
225					230					235					240
Gly	Val	Val	Ala	Ala	Asn	Ile	Ser	Asn	Tyr	Asn	Ala	Ser	Ser	Arg	Ile
				245					250					255	
Asp	Val	Asp	Gly	Tyr	Leu	His	Ala	Lys	Lys	Arg	Leu	Asn	Val	Glu	Ala
			260					265					270		
His	Asn	Ile	Thr	Lys	Asn	Ser	Val	Leu	Gln	Thr	Gly	Ser	Asp	Leu	Gly
		275					280					285			
Thr	Ser	Lys	Phe	Met	Asn	Asp	His	Val	Tyr	Glu	Ser	Gly	His	Leu	Lys
	290					295					300				
Ser	Ile	Leu	Asp	Ala	Ile	Lys	Gln	Arg	Phe	Gly	Gly	Asp	Ser	Val	Asn
305					310					315					320
Glu	Glu	Ile	Lys	Asn	Lys	Leu	Thr	Asn	Leu	Phe	Ser	Val	Gly	Val	Ser
				325					330					335	
Ala	Thr	Ile	Ala	Asn	His	Asn	Asn	Ser	Ala	Ser	Val	Ala	Ile	Gly	Glu
			340					345					350		
Ser	Gly	Arg	Leu	Ser	Ser	Gly	Val	Glu	Gly	Ser	Asn	Val	Arg	Ala	Leu
		355					360					365			
Asn	Glu	Ala	Gln	Asn	Leu	Arg	Ala	Thr	Thr	Ser	Ser	Gly	Ser	Val	Ala
	370					375					380				
Val	Arg	Lys	Glu	Glu	Lys	Lys	Lys	Leu	Ile	Gly	Asn	Ala	Ala	Val	Phe
385					390					395					400
Tyr	Gly	Asn	Tyr	Lys	Asn	Asn	Ala	Ser	Val	Thr	Ile	Ala	Asp	His	Ala
				405					410					415	
Glu	Leu	Val	Ser	Glu	Gly	Lys	Ile	Asp	Ile	Asn	Ser	Glu	Asn	Lys	Ile
			420					425					430		
Glu	Tyr	Lys	Asn	Pro	Ser	Lys	Met	Ala	Lys	Ser	Val	Ile	Asp	Lys	Leu
		435					440					445			
Glu	Leu	Leu	Lys	Arg	Ala	Phe	Gly	Lys	Glu	Thr	Lys	Thr	Pro	Glu	Tyr
	450					455					460				
Asp	Pro	Lys	Asp	Ile	Glu	Ser	Ile	Glu	Lys	Leu	Leu	Asn	Ala	Phe	Ser
465					470					475					480
Glu	Lys	Leu	Asp	Gly	Lys	Pro	Glu	Leu	Leu	Leu	Asn	Gly	Glu	Arg	Met
				485					490					495	
Thr	Ile	Ile	Leu	Pro	Asp	Gly	Thr	Ser	Lys	Thr	Gly	Thr	Ala	Ile	Glu
			500					505					510		
Ile	Ala	Asn	Tyr	Val	Gln	Gly	Glu	Met	Lys	Lys	Leu	Glu	Glu	Lys	Leu
		515					520					525			
Pro	Lys	Gly	Phe	Lys	Ala	Phe	Ser	Glu	Gly	Leu	Ser	Gly	Leu	Ile	Lys
	530					535					540				
Glu	Thr	Leu	Asn	Phe	Thr	Gly	Val	Gly	Asn	Tyr	Ala	Asn	Phe	His	Thr
545					550					555					560

PCT302~1

Phe Thr Ser Ser Gly Ala Asn Gly Glu Arg Asp Val Ser Ser Val Gly
 565 570 575
 Gly Ala Val Ser Trp Val Glu Gln Glu Asn Tyr Ser Lys Val Ser Val
 580 585 590
 Gly Lys Gly Ala Lys Leu Ala Ala Lys Lys Asp Leu Asn Ile Lys Ala
 595 600 605
 Ile Asn Lys Ala Glu Thr Val Asn Leu Val Gly Asn Ile Gly Leu Ala
 610 615 620
 Arg Ser Ser Thr Ser Gly Ser Ala Val Gly Gly Arg Leu Asn Val Gln
 625 630 635 640
 Arg Ser Lys Asn Ser Ala Ile Val Glu Ala Lys Glu Lys Ala Glu Leu
 645 650 655
 Ser Gly Glu Asn Ile Asn Ala Asp Ala Leu Asn Arg Leu Phe His Val
 660 665 670
 Ala Gly Ser Phe Asn Gly Gly Ser Gly Gly Asn Ala Ile Asn Gly Met
 675 680 685
 Gly Ser Tyr Ser Gly Gly Ile Ser Lys Ala Arg Val Ser Ile Asp Asp
 690 695 700
 Glu Ala Tyr Leu Lys Ala Asn Lys Lys Ile Ala Leu Asn Ser Lys Asn
 705 710 715 720
 Asp Thr Ser Val Trp Asn Ala Ala Gly Ser Ala Gly Ile Gly Thr Lys
 725 730 735
 Asn Ala Ala Val Gly Val Ala Val Ala Val Asn Asp Tyr Asp Ile Ser
 740 745 750
 Asn Lys Ala Ser Ile Glu Asp Asn Asp Glu Gly Gln Ser Lys Tyr Asp
 755 760 765
 Lys Asn Lys Asp Asp Glu Val Thr Val Thr Ala Glu Ser Leu Glu Val
 770 775 780
 Asp Ala Lys Thr Thr Gly Thr Ile Asn Ser Ile Ser Val Ala Gly Gly
 785 790 795 800
 Ile Asn Lys Val Gly Ser Lys Pro Ser Glu Glu Lys Pro Lys Ser Glu
 805 810 815
 Glu Arg Pro Glu Gly Phe Phe Gly Lys Ile Gly Asn Lys Val Asp Ser
 820 825 830
 Val Lys Asn Lys Ile Thr Asp Ser Met Asp Ser Leu Thr Glu Lys Ile
 835 840 845
 Thr Asn Tyr Ile Ser Glu Gly Val Lys Lys Ala Gly Asn Leu Pro Ser
 850 855 860
 Asn Val Ser His Thr Pro Asp Lys Gly Pro Ser Phe Ser Leu Gly Ala
 865 870 875 880
 Ser Gly Ser Val Ser Phe Asn Asn Ile Lys Lys Glu Thr Ser Ala Val
 885 890 895
 Val Asp Gly Val Lys Ile Asn Leu Lys Gly Ala Asn Lys Lys Val Glu
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 Val Thr Ser Ser Asp Ser Thr Phe Val Gly Ala Trp Gly Gly Ser

915 920 925
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 <211> 580
 <212> PRT
 <213> *Fusobacterium necrophorum*

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 Asp Gly Val Lys Asn Lys Gly Ala Asn Lys Lys Val Val Thr Ser Ser
 20 25 30
 Asp Ser Thr Val Gly Ala Trp Gly Gly Ser Ala Ala Trp Asn His Gly
 35 40 45
 Ser Gly Asn Ser Asn Ser Ala Gly Ala Gly Ala Ala Val Asn Asn
 50 55 60
 Ser Lys Thr Ser Ala Val Lys Asn Ser Asp Arg Asn Ala Asn Lys Lys
 65 70 75 80
 Val Asn Ala Ser Gly Gly Thr Val Ala Ala Gly Ala Gly Ala Val Lys
 85 90 95
 Ser Gly Gly Gly Lys Ser Tyr Gly Thr Ser Ala Ser Asn Val Asn Asn
 100 105 110
 Val Ser Ala Lys Ser Asn Asn Thr Val Ala Gly Ser Ser Lys Met Asp
 115 120 125
 Val Asp Val Thr Ala Tyr Ala Asp Thr Val Thr Gly Ala Asn Ala Gly
 130 135 140
 Lys Ser Asn Gly Thr Val Gly Ala Thr Val Thr Val Ala Lys Asn Asn
 145 150 155 160
 Lys Val Asn Ala Ser Ser Gly Gly Arg Tyr Thr Asn Val Asn Arg Ala
 165 170 175
 Asp Ala Lys Ala Ala Thr Thr Val Thr Ala Ala Val Thr Thr Gly Gly
 180 185 190
 Thr Ser Ser Gly Ala Gly Gly Asn Tyr Gly Ala Val Ser Val Asn Lys
 195 200 205
 Asp Asn Asp Val Ala Ser Val Asp Lys Ser Ser Gly Ala Asn Asn Val
 210 215 220
 Ala Lys Asp Val Lys Gly Ser Ser Asp Ala Lys Tyr Ala Asn Gly Lys
 225 230 235 240
 Asp Lys Lys Tyr Asp Arg Gly Asn Thr Thr Gly Asn Gly Tyr Tyr Thr
 245 250 255
 Lys Lys Ala Lys Lys Lys Gly Ala Val Val Asn Ala Ala Ser Val Ala
 260 265 270
 Gly Thr Asp Lys Ser Ala Gly Gly Val Ala Ala Val Asn Thr Val Lys
 275 280 285

PCT302~1

Asn Lys Lys Ala Ser Gly Ser Asn Lys Ala Gly Asp Lys His Ala Lys
 290 295 300
 His Val Asn Val Ala Lys Ser Ser Thr Val Val Val Asn Ala Ala Ser
 305 310 315 320
 Gly Ala Ser Lys Asp Ala Ser Gly Met Gly Ser Gly Ala Trp Asp Ser
 325 330 335
 Asn Asp Thr Ala Lys Val Asp Lys Gly Arg Ser Ala Asp Ser Asn Val
 340 345 350
 Asn Ala Asn Asn Ser Gly Val Asn Val Ala Gly Thr Ala Gly Ser Ser
 355 360 365
 Thr Ala Val Gly Ala Ala Ala Asn Asn Thr His Asn Lys Thr Ser Ala
 370 375 380
 Thr Gly Thr Lys Val Asn Ser Gly Lys Asn Thr Lys Val Asn Val Ala
 385 390 395 400
 Asn Asp Ser His Thr Asn Val Ser Ala Gly Gly Ala Ala Ser Lys Ala
 405 410 415
 Gly Gly Gly Met Val Ser Val Asn Arg Gly Ser Asp Thr Ala Val Ser
 420 425 430
 Asp Ser Gly Val Ser Ser Asn Val Asp Ala Lys Asp Lys Thr Asn Thr
 435 440 445
 Ala Gly Asn Ala Asn Gly Gly Lys Ala Ala Gly Val Gly Ala Thr Val
 450 455 460
 Ala His Thr Asn Gly Lys Ser Val Ala Val Lys Asn Ser Lys Thr Thr
 465 470 475 480
 Ala Asn Asp Asp Arg Lys Asn Asn Val Thr Ala Lys Asp Tyr Thr Met
 485 490 495
 Thr Asn Thr Ala Val Gly Val Gly Gly Ala Lys Gly Ala Ser Val Gly
 500 505 510
 Ala Ser Ala Ser Thr Thr Asn Lys Thr Val Ser Ser His Val Asp Thr
 515 520 525
 Asp Asp Lys Asp Asn Asn Gly Asn Lys Lys Ala Asn Val Asn Val Ala
 530 535 540
 Asn Thr Ser Val Val Thr Asn Ala Thr Val Ser Gly Ala Ser Gly Ala
 545 550 555 560
 Ala Val Gly Ala Gly Val Ala Val Asn Lys Thr Asn Thr Ser Ala His
 565 570 575
 Lys Asn Ser Thr
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<210> 5

<211> 628

<212> PRT

<213> Fusobacterium necrophorum

<400> 5

PCT302~1

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Ala His Ile Lys Asn Ser Thr Gln Asn Val Arg Asn Ala Leu Val Lys
20 25 30
Ser Lys Ser His Ser Ser Ile Lys Thr Ile Gly Ile Gly Ala Gly Val
35 40 45
Gly Ala Gly Gly Ala Gly Val Thr Gly Ser Val Ala Val Asn Lys Ile
50 55 60
Val Asn Asn Thr Ile Ala Glu Leu Asn His Ala Lys Ile Thr Ala Lys
65 70 75 80
Gly Asn Val Gly Val Ile Thr Glu Ser Asp Ala Val Ile Ala Asn Tyr
85 90 95
Ala Gly Thr Val Ser Gly Val Ala Arg Ala Ala Ile Gly Ala Ser Thr
100 105 110
Ser Val Asn Glu Ile Thr Gly Ser Thr Lys Ala Tyr Val Lys Asp Ser
115 120 125
Thr Val Ile Ala Lys Glu Glu Thr Asp Asp Tyr Ile Thr Thr Gln Gly
130 135 140
Gln Val Asp Lys Val Val Asp Lys Val Phe Lys Asn Leu Asn Ile Asn
145 150 155 160
Glu Asp Leu Ser Gln Lys Arg Lys Ile Ser Asn Lys Lys Gly Phe Val
165 170 175
Thr Asn Ser Ser Ala Thr His Thr Leu Lys Ser Leu Leu Ala Asn Ala
180 185 190
Ala Gly Ser Gly Gln Ala Gly Val Ala Gly Thr Val Asn Ile Asn Lys
195 200 205
Val Tyr Gly Glu Thr Glu Ala Leu Val Glu Asn Ser Ile Leu Asn Ala
210 215 220
Lys His Tyr Ser Val Lys Ser Gly Asp Tyr Thr Asn Ser Ile Gly Val
225 230 235 240
Val Gly Ser Val Gly Val Gly Gly Asn Val Gly Val Gly Ala Ser Ser
245 250 255
Asp Thr Asn Ile Ile Lys Arg Asn Thr Lys Thr Arg Val Gly Lys Thr
260 265 270
Thr Met Ser Asp Glu Gly Phe Gly Glu Glu Ala Glu Ile Thr Ala Asp
275 280 285
Ser Lys Gln Gly Ile Ser Ser Phe Gly Val Gly Val Ala Ala Ala Gly
290 295 300
Val Gly Ala Gly Val Ala Gly Thr Val Ser Val Asn Gln Phe Ala Gly
305 310 315 320
Lys Thr Glu Val Asp Val Glu Glu Ala Lys Ile Leu Val Lys Lys Ala
325 330 335
Glu Ile Thr Ala Lys Arg Tyr Ser Ser Val Ala Ile Gly Asn Ala Ala
340 345 350
Val Gly Val Ala Ala Lys Gly Ala Gly Ile Gly Ala Ala Val Ala Val

PCT302~1

355 360 365
 Thr Lys Asp Glu Ser Asn Thr Arg Ala Arg Val Lys Asn Ser Lys Ile
 370 375 380
 Met Thr Arg Asn Lys Leu Asp Val Ile Ala Glu Asn Glu Ile Lys Ser
 385 390 395 400
 Gly Thr Gly Ile Gly Ser Ala Gly Ala Gly Ile Leu Ala Ala Gly Val
 405 410 415
 Ser Gly Val Val Ser Val Asn Asn Ile Ala Asn Lys Val Glu Thr Asp
 420 425 430
 Ile Asp His Ser Thr Leu His Ser Ser Thr Asp Val Asn Val Lys Ala
 435 440 445
 Leu Asn Lys Ile Ser Asn Ser Leu Thr Ala Gly Gly Gly Ala Ala Gly
 450 455 460
 Leu Ala Ala Val Thr Gly Val Val Ser Val Asn Thr Ile Asn Ser Ser
 465 470 475 480
 Val Ile Ala Arg Val His Asn Asn Ser Asp Leu Thr Ser Val Arg Glu
 485 490 495
 Lys Val Asn Val Thr Ala Lys Glu Glu Lys Asn Ile Lys Gln Thr Ala
 500 505 510
 Ala Asn Ala Gly Ile Gly Gly Ala Ala Ile Gly Ala Asn Val Leu Val
 515 520 525
 Asn Asn Phe Gly Thr Ala Val Glu Asp Arg Lys Asn Ser Glu Gly Lys
 530 535 540
 Gly Thr Glu Val Leu Lys Thr Leu Asp Glu Val Asn Lys Glu Gln Asp
 545 550 555 560
 Lys Lys Val Asn Asp Ala Thr Lys Lys Ile Leu Gln Ser Ala Gly Ile
 565 570 575
 Ser Thr Glu Asp Thr Ser Val Lys Ala Asp Arg Gly Asp Thr Gln Gly
 580 585 590
 Glu Gly Ile Lys Ala Ile Val Lys Thr Ser Asp Ile Ile Gly Lys Asn
 595 600 605
 Val Asp Ile Thr Thr Glu Asp Lys Asn Asn Ile Thr Ser Thr Gly Gly
 610 615 620
 Leu Gly Thr Ala
 625
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 <211> 773
 <212> PRT
 <213> *Fusobacterium necrophorum*

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PCT302~1

Asp Ile Thr Thr Glu Asp Lys Asn Asn Ile Thr Ser Thr Gly Gly Leu
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 Gly Thr Ala Gly Leu Ala Ser Ala Ser Gly Thr Val Ala Val Thr Asn
 35 40 45
 Ile Lys Arg Asn Ser Gly Val Thr Val Glu Asn Ser Phe Val Lys Ala
 50 55 60
 Ala Glu Lys Val Asn Val Arg Ser Asp Ile Thr Gly Asn Val Ala Leu
 65 70 75 80
 Thr Ala Tyr Gln Gly Pro Val Gly Ala Leu Gly Ile Gly Ala Ala Tyr
 85 90 95
 Ala Glu Leu Asn Ser Asn Gly Arg Ser Asn Ile Ser Ile Lys Asn Ser
 100 105 110
 Lys Leu Leu Gly Lys Asn Ile Asp Val Ile Val Lys Asp Lys Ser Glu
 115 120 125
 Leu Arg Ala Glu Ala Lys Gly Leu Thr Val Gly Ala Val Ala Ala Gly
 130 135 140
 Ala Ile Ile Ser Lys Ala Lys Asn Glu Met Asn Ser Glu Val Glu Ile
 145 150 155 160
 Glu Lys Ser Ile Phe Asn Glu Glu Asn Arg Val Thr Ser Pro Ser Lys
 165 170 175
 Gly Ile Gly Arg Glu Ile Asn Val Lys Val Glu Lys Glu Asn Arg Val
 180 185 190
 Thr Ala Glu Ser Gln Gly Ala Ser Val Gly Ala Val Ala Gly Ala Gly
 195 200 205
 Ile Ile Ser Glu Ala Lys Asp Ala Gly Ser Ser Tyr Leu Lys Val Ser
 210 215 220
 Thr Lys Ser Gly Arg Ser Ile Phe His Ala Asp Asn Val Asn Met Glu
 225 230 235 240
 Ala Thr His Lys Met Lys Val Thr Ala Val Ser Lys Ala Val Thr Gly
 245 250 255
 Ser Val Leu Gly Gly Val Gly Val Thr Lys Ala Glu Ala Thr Ala Ala
 260 265 270
 Gly Lys Thr Met Val Glu Val Glu Glu Gly Asn Leu Phe Arg Thr Asn
 275 280 285
 Arg Leu Asn Ala Ile Ser Lys Val Glu Gly Leu Asp Glu Asp Lys Val
 290 295 300
 Thr Ala Lys Ser Ser Val Val Ser Gly Asn Gly Gly Gly Ile Ala Gly
 305 310 315 320
 Ala Gly Val Asn Thr Ser Thr Ala Gln Ser Asn Thr Glu Ser Val Val
 325 330 335
 Arg Leu Arg Lys Gln Asp Tyr Glu Asn Asn Asp Tyr Thr Lys Lys Tyr
 340 345 350
 Ile Ser Glu Val Asn Ala Leu Ala Leu Asn Asp Thr Lys Asn Glu Ala
 355 360 365
 Asn Ile Glu Ser Leu Ala Val Ala Gly Val His Ala Gln Gly Thr Asn
 370 375 380

PCT302~1

Lys Ala Phe Thr Arg Ser Asn Lys Leu Thr Ser Thr Thr Val Asn Gly
 385 390 395 400
 Gly Asn Val Ser Gln Leu Arg Ala Lys Ala Leu Ala Lys Asn Glu Asn
 405 410 415
 Tyr Gly Asn Val Lys Gly Thr Gly Gly Ala Leu Val Gly Ala Glu Thr
 420 425 430
 Ala Ala Val Glu Asn Tyr Thr Lys Ser Thr Thr Gly Ala Leu Val Ala
 435 440 445
 Gly Asn Trp Glu Ile Gly Asp Lys Leu Glu Thr Ile Ala Arg Asp Asn
 450 455 460
 Thr Ile Val Arg Val Asn Gly Asp Gly Thr Lys Gly Gly Leu Val Gly
 465 470 475 480
 Lys Asn Gly Ile Ser Val Lys Asn Thr Ile Ser Gly Glu Thr Lys Ser
 485 490 495
 Ser Ile Glu Asp Lys Ala Arg Ile Val Gly Thr Gly Ser Val Asn Val
 500 505 510
 Asp Ala Leu Asn Glu Leu Asp Val Asp Leu Gln Gly Lys Ser Gly Gly
 515 520 525
 Tyr Gly Gly Ile Gly Ile Gly Asn Val Asp Val Asn Asn Val Ile Lys
 530 535 540
 Lys Asn Val Glu Ala Lys Ile Gly Arg His Ala Ile Val Glu Thr Thr
 545 550 555 560
 Gly Lys Gln Glu Tyr Gln Ala Phe Thr Arg Ala Lys Val Asn Ile Leu
 565 570 575
 Gly Lys Gly Asp Ala Ala Ala Ala Ala Ile Ser Asn Val His Ile
 580 585 590
 Ser Asn Glu Met Asp Ile Lys Asn Leu Ala Lys Gln Tyr Ala Ser Ser
 595 600 605
 Gln Leu Ile Thr Lys Asn Ser Lys Asn Asn Ile Thr Leu Ala Ser Ser
 610 615 620
 Ser Glu Ser Asn Val Asn Val His Gly Val Ala Glu Ala Arg Gly Ala
 625 630 635 640
 Gly Ala Lys Ala Thr Val Ser Val Lys Asn Gln Ile Asn Arg Thr Asn
 645 650 655
 Asn Val Asp Leu Ala Gly Lys Ile Lys Thr Glu Gly Asn Ile Asn Val
 660 665 670
 Tyr Ala Gly Tyr Asp Lys Asn Tyr Asn Ile Ser Lys Thr Asn Ser Lys
 675 680 685
 Ala Ile Ala Asp Ala Lys Ser His Ala Ala Ala Ala Ser Ala Thr Ala
 690 695 700
 Thr Ile Glu Lys Asn Glu Val Lys Phe Asn Asn Ala Ile Arg Glu Phe
 705 710 715 720
 Lys Asn Asn Leu Ala Arg Leu Glu Gly Lys Ala Asn Lys Lys Thr Ser
 725 730 735
 Val Gly Ser Asn Gln Val Asp Trp Tyr Thr Asp Lys Tyr Thr Trp His

PCT302~1

740 745 750
 Ser Ser Glu Lys Ala Tyr Lys Lys Leu Thr Tyr Gln Ser Lys Arg Gly
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 Glu Lys Gly Lys Lys
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 <211> 338
 <212> PRT
 <213> Fusobacterium necrophorum

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 Leu Gly Gln Lys Asn Thr Gly Glu Leu Arg Trp Gly Leu Asn Tyr Ile
 35 40 45
 Asn Asn Ser Val Thr Gly Asn Arg Asp Gln Leu Ser Leu Thr Ser Leu
 50 55 60
 Val Thr Glu Gly Thr Ala Ser Leu Ser Ser Phe Tyr Thr Phe Pro Val
 65 70 75 80
 Ser Lys Lys Gly Thr Lys Ile Ser Leu Gln His Ser Val Gly Lys Leu
 85 90 95
 Lys His Ile Gln Gly Ala Leu Lys His Lys Ile Thr Gly Asn Ser Tyr
 100 105 110
 Ser Tyr Gly Val Gly Ile Val His Pro Ile Leu Val His Glu Lys Asn
 115 120 125
 Lys Val Glu Leu Ser Leu Asp Trp Val Lys Gln Arg Thr Val Thr Asp
 130 135 140
 Leu Leu Lys Leu Lys Trp Val Asn Asn Arg Leu Ser Lys Tyr Thr Ala
 145 150 155 160
 Gly Ile Gly Ile Ser His Tyr Glu Glu Asp Ser Val Phe Tyr Thr Lys
 165 170 175
 Gln Asn Ile Thr Lys Gly Lys Phe Ile Pro Ile Ser Gly Asp Ala Arg
 180 185 190
 Asn Tyr Thr Lys Tyr Asp Met Phe Leu Ile Tyr Gln Lys Asn Leu Lys
 195 200 205
 Tyr Asn Thr Leu Val Thr Leu Lys Met Ala Gly Gln Tyr Ser Leu Ser
 210 215 220
 Lys Lys Leu Pro Ser Val Glu Gln Ile Tyr Ala Gly Gly Ala Tyr Asn
 225 230 235 240
 Val Arg Gly Tyr Pro Glu Asn Phe Met Gly Ala Glu His Gly Val Phe
 245 250 255

PCT302~1

Phe Asn Ala Glu Leu Ser Lys Leu Val Glu Asn Lys Gly Glu Phe Phe
 260 265 270

Val Phe Leu Asp Gly Ala Ser Leu His Gly Glu Ser Ala Trp Gln Glu
 275 280 285

Asn Arg Ile Phe Ser Ser Gly Phe Gly Tyr Lys Ile Arg Phe Leu Glu
 290 295 300

Lys Asn Asn Ile Ala Val Ser Met Ala Phe Pro Trp Lys Lys Lys Ile
 305 310 315 320

Asn Ser Ile Ser Val Asp Ser Asn Arg Ile Tyr Ile Thr Ile Asn His
 325 330 335

Glu Phe

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 <211> 9726
 <212> DNA
 <213> Fusobacterium necrophorum

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 accgcgactg agaatttttg aacaaaaata gaaaaaaagg ataatgttta tgacattact 180
 acaaacaaga ttcaagggga gaacgctttt aacagtttta atagatttgc tttaacagaa 240
 aataatatag caaatctata ttttggggaa aagaatagta cgggggtaaa taatcttttt 300
 aactttgtca atggaaaaat tgaagtagat gggattatca acggaattcg agaaaataaa 360
 attggaggaa atttatattt cttaagctcg gaagggatgg cagtaggaaa aaatggagtt 420
 atcaatgctg gttcttttca ttctattatt ccaaaacaag atgattttta gaaggctttg 480
 gaagaagcca aacatggtaa agtttttaat ggaatcattc cagtagatgg aaaagtaaaa 540
 attccattga atccgaatgg aagcattacg gtagaaggaa aaatcaatgc tgttgaaggc 600
 atcggtttat atgcggcgga tattagattg aaagatactg caatactaaa gacaggaatt 660
 acagatttta aaaatttagt caatattagt gatcgaataa attctggtct gaccggagat 720
 ttaaaagcta ccaagacaaa atctggagat attattcttt cagctcacat agattctcct 780
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<210> 11

<211> 2141

<212> DNA

<213> *Fusobacterium necrophorum*

<400> 11

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<210> 12

<211> 1887

<212> DNA

<213> *Fusobacterium necrophorum*

<400> 12

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aaacaattgg	aattggagct	ggagttggag	ctggaggagc	tggagtgaca	ggttctgtag	180

PCT302~1

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<210> 13

<211> 2322

<212> DNA

<213> *Fusobacterium necrophorum*

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<210> 14

<211> 1017

<212> DNA

<213> *Fusobacterium necrophorum*

<400> 14

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<210> 15

<211> 11130

<212> DNA

<213> *Fusobacterium necrophorum*

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